

Fig. 6A

C.anc.env (subtype C ancestral env. The amino acid sequence is different from Los Alamos Database August 2002)

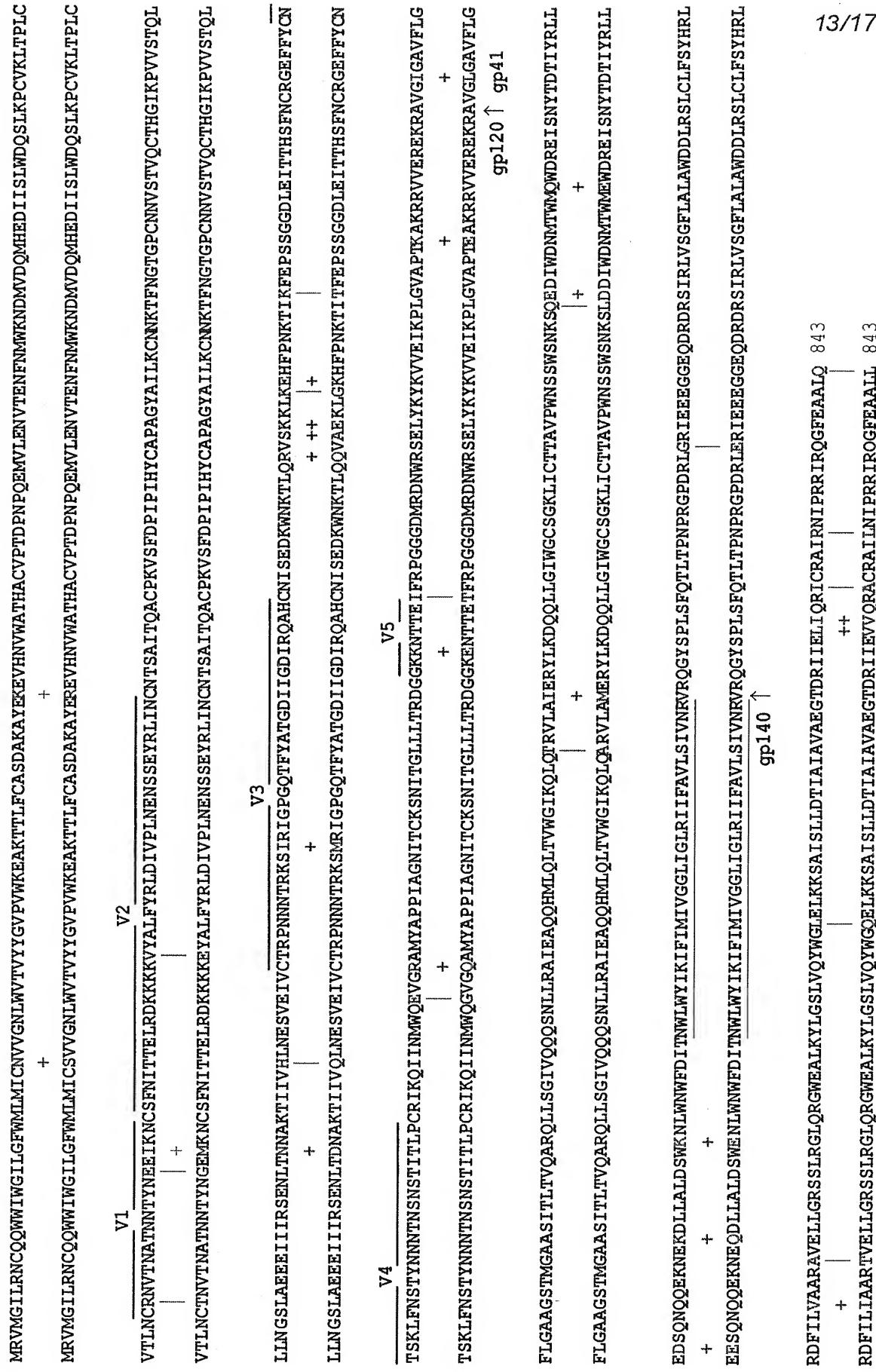
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CATCGAGGTGGTGCAGCGCGCTGCGCCGCATCCTGAACATCCCCCGCC
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Fig. 6B

C.con.env (subtype C consensus env. The amino acid sequence is different from Los Alamos Database August 2002)

GGCGCCATGCGCGTGTGGGCATCCTGCGCAACTGCCAGCAGTGGTGGAT
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Fig. 8



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Fig. 14B

CONs.env (gorup M consensus env gene. This one contain the consensus sequence for variable regions in env gene. The identical amino acid sequences as in the public domain)

```
CCGCCGCCATGCGCGTGCAGCGCATCCAGCGAACTGCCAGCACCTGTG  
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Fig. 15A

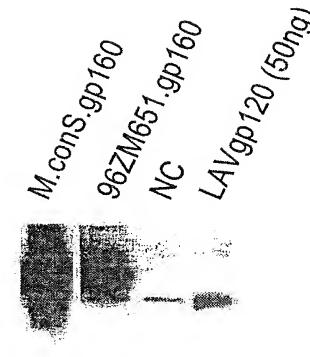
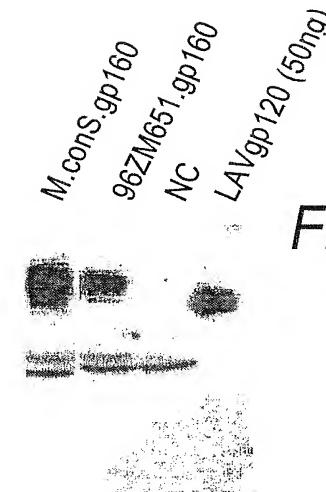


Fig. 15B

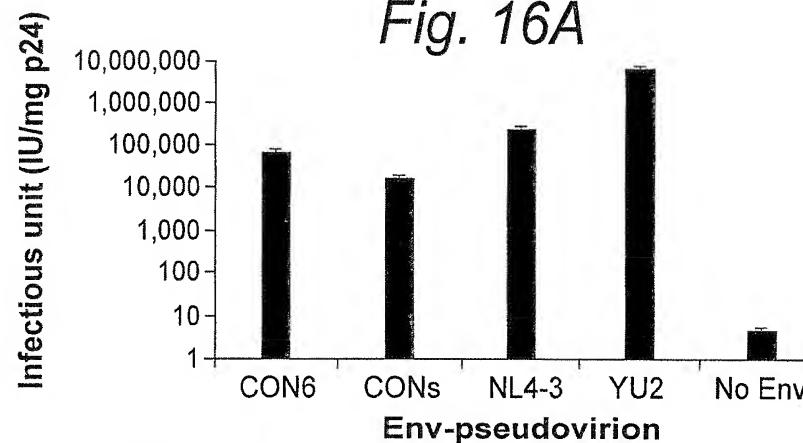


Cell lysate

Supernatant

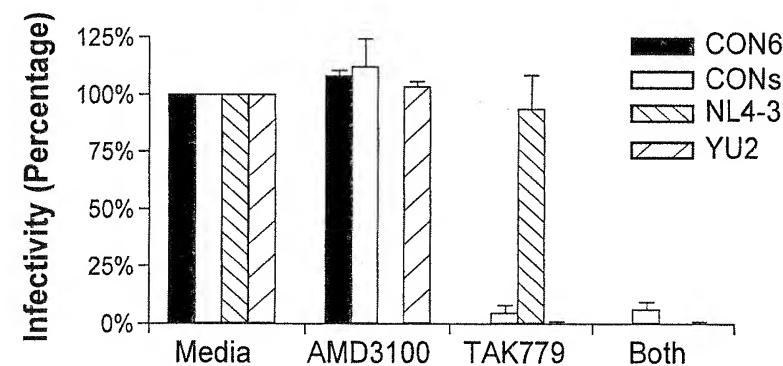
Expression of A.con env gene in mammalian cells

Fig. 16A



Infectivity and coreceptor usage of CON6 and CONs env genes

Fig. 16B



Infectivity and coreceptor usage of CON6 and CONs env genes

Fig. 18B

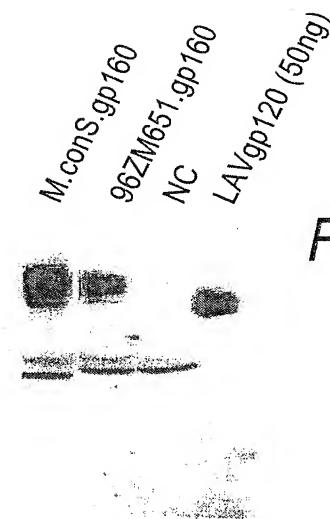
A.con.env (subtype A consensus env. Identical amino acid sequence to that in the public domain)

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TGGAACCTGCTGCTGACTGGGGCCGAGCTGAAGATCTCCGCCATCAA
CCTGCTGGACACCACATGCCATGCCGTGGCGGCTGGACCGACCGCGTGA
TCGAGATCGGCCAGCGCATCTGCCGCCATCCTGAACACATCCCCGCC
ATCCGCCAGGGCCTGGAGCGCGCCCTGCTGTAA

Fig. 18C



Fig. 18D



Cell lysate

Supernatant

Expression of A.con env gene in mammalian cells

M.con.gag (group M consensus gag. Identical amino acid sequence to that in the public domain)

CCGCCGCCATGGCGCCGCCCTCCGTGCTGTCCGGGGCAAGCTGGA
CGCCTGGAGAAGATCCGCCTGCGCCCCGGCGCAAGAAGAAGTACCGCC
TGAAGCACCTGGTGTGGGCCTCCCGCAGCTGGAGGGCTGCAAGCAGATCATCGGCCAGCT
CCCGGCCTGCTGGAGACCTCCGAGGGCTGCAAGCAGATCATCGGCCAGCT
GCAGCCCCGCCCTGAGACCGGCTCCGAGGAGCTGCGCTCCGTACAACA
CCGTGGCCACCCCTGTACTGCGTGCACCAGCGCATCGAGGTGAAGGACACC
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TAA

Fig. 19A

M.con.pol.nuc

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Fig. 19B

Fig. 19C

M.con.nef (group M consensus nef. Identical amino acid sequence to that in the public domain)

```
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GGTGTACCAACCCAGGGCTACTTCCCGAATGGCAGAAGACTACACCCCG  
GCCCGGCACTCGCTACCCCCTGACCTCGCTGGTGCTTCAAGCTGGTG  
CCCGTGGACCCGAGGAGGTGGAGGAGGCCAACGAGGGCGAGAACAACTC  
CCTGCTGCAACCCATGTGCCAGCACGGCATGGAGGAGGAAGCGAGG  
TGCTGATGTGGAAAGTTCGACTCCCGCCTGGCCCAATTCGCCCC  
GAGCTGCACCCGAGTAACAAGGACTGCTAA
```

Fig. 19D

C.con.pol.nuc

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CCGCGGCCATGGCCAGATCACCTGTGGCAGCGCCCCCTGGTGTCCAT  
CAAGGTGGCGGCCAGATCAAGGAGGCCATGCTGGCCACCGCGCCGACG  
ACACCGTGCTGGAGGAGATCAAACCTGCCGGCAAGTGGAAAGCCCAAGATG  
ATCGCGGCCATGGCGCTTCATCAAGGTGCGCCAGTACGACCAAGATCCT  
GATCGAGATCTGCGCAAGAAGGCCATCGCACCGTGTGGTGGGCCCA  
CCCCCGTGAACATCATCGGCCAACATGCTGACCCAGCTGGCTGCACC  
CTGAACCTCCCCATCTCCCCATCGAGACCGTGTGGCTGAAGCTGAAGCC  
CGGCATGGACGGCCCAAGGTGAAGCAGTGGCCCTGACCGAGGAGAAGA  
TCAAGGCCCTGACCGCCATCTGCGAGGAGATGGAGAAGGAGGGCAAGATC  
ACCAAGATCGGCCCCGAGAACCCCTACAACACCCCCGTGTCGCCATCAA  
GAAGAAGGACTCCACCAAGTGGCGCAAGCTGGGACTTCCGCGAGCTGA  
ACAAGCGACCCAGGACTCTGGAGGTGCAGCTGGCATCCCCCACCC  
GCCGGCTGAAGAAGAAGTCCGTGACCGTGTGGACGTGGCGACGC  
CTACTTCTCCGTGCCCTGGACGAGGGCTCCGCAAGTACACCGCCTTCA  
CCATCCCCCTCCATCAACAACGAGACCCCCGGCATCCGCTACCAAGTACAAC  
GTGCTGCCCAAGGGCTGGAAGGGCTCCCCCGCCATCTCCAGTCCCTCAT  
GACCAAGATCTGGAGCCCTTCCGCGCCAGAACCCCGAGATCGTGTGATCT  
ACCAAGTACATGGACGACCTGTACGTGGCTCCGACCTGGAGATCGGCCAG  
CACCGCGCCAAGATCGAGGAGCTGCGCGAGCACCTGCTGAAGTGGGCTT  
CACCACCCCGACAAGAAGCACCAGAAGGAGCCCCCTCCTGTGGATGG  
GCTACGAGCTGCACCCGACAAGTGGACCGTGCAGCCCATCCAGCTGCC  
GAGAAGGACTCCTGGACCGTGAACGACATCCAGAAGCTGGTGGCAAGCT  
GAACCTGGCCTCCCAGATCTACCCCGCATCAAGGTGCCAGCTGTGCA  
AGCTGCTGCGCGGCCAAGGCCCTGACCGACATCGTGCCCTGACCGAG  
GAGGCCGAGCTGGAGCTGGCCGAGAACCGCGAGATCCTGAAGGAGGCCGT  
GCACGGCGTGTACTACGACCCCTCAAGGACCTGATCGCCGAGATCCAGA  
AGCAGGGCCACGACCAGTGGACCTACCAAGATCTACCAAGGAGGCCCTCAAG  
AACCTCAAGACCGCAAGTACGCCAAGATGCGCACCGCCACACCAACGA  
CGTGAAGCAGCTGACCGAGGCCGTGAGAAGATCGCCATGGAGTCCATCG  
TGATCTGGGCAAGACCCCCCAAGTTCCGCTGCCATCCAGAAGGAGAC  
TGGGAGACCTGGGACCGACTACTGGCAGGCCACCTGGATTCCGAGTG  
GGAGTTCGTGAACACCCCCCTGGTGAAGCTGTGGTACCAAGCTGGAGA  
AGGAGCCCATCGCCGGCGAGACCTTACGTGGACGGCGCCGCAAC
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CGCGAGACCAAGATCAGGCCAAGGCCGCTAACGTGACCGAACCAACCAAGAAAACCGAGCTGCAGG
GAAGATCGTGTCCCTGACCGAGACCAACCAAGAAAACCGAGCTGCAGG
CCATCCAGCTGGCCTGCAAGGACTCCGGCTCCAGGTGAACATCGTGACC
GAATCCAGTACGCCCTGGCATCATCCAGGCCAGCCGACAAGTCCGA
GTCCGAGCTGGTGAACCATCATCGAGCTGATCAAGAAGGAGCG
TGTACCTGTCTGGGTGCCAGATCAAGGGCATCGGGCAACGGAGCAG
GTGGACAAAGCTGGTCTCGGGCATCGCAAGGTGCTGTTCTGGACGG
CATCGACAAAGGCCAGGAGGAGGAGGAGCTAACACTCCAACTGGCCGC
CCATGGCCTCCGAGTTCAACCTGCCCTCCCATCGTGGCAAGGAGATCGTG
GCCTCTGGACAAAGTGCAGCTGAAGGGCCATGCCAGGAGGGCCAGGT
GGACTGCTCCCCCGCATCGCAGCTGGAGCTGGACTGACCCACCTGGAGGGCA
AGATCATCTGGTGGCGTGCACGTGGCTCCGGCTACATCGAGGCCAG
GTGATCCCCGGAGACCGGCCAGGGAGACCCGCTACTCATCTGAAGGCT
GCCGGCGCTGGCCCGTGAAGGTGATCCACACCGACAAAGGCTCCAAC
TCACCTCGCCGCGCTGAAGGCCGCTGCTGGTGGCCGATCCAGCGAG
GAGTTGGCATCCCTACAACCCCCAGTCCCAGGGCTGGGTGGAGTCCAT
GAACAAGGAGCTGAAGAAGATCATGGCCAGGTGGCGGACCCAGGGCGAGC
ACCTCAAGGACCCGGCTGCAATGGCCGTTGTCATCCACAACTTCAAGCGC
AAGGGGGCATCGGGCTACTCCGGCATCGACATCATCGACATCATCGACAT
CGCCACCGACATCCAGACCCAGGAGCTGCAGAAGCAGATCATCAAGATCC
AGAACTTCCGGTGTACTACCGCGACTCCCCGGACCCCATCTGGAAAGGGC
CCGCCAAGCTGCTGTGAAGGGGAGGGGGCCAGGGCAAGGCCAAGGACT
CTCCGACATCAAGGTGGTGGCCGGCAAGGCCAAGGACTCATCAAGGACT
ACGGCAAGCAGATGGCCGGCGCCGACTCGTGGCCGGCAGGAGCAG
GACTAA

Fig. 19D (continued)

M.con.gag (group M consensus gag)

MGARASVLSGGKIDAEWEKIRLRPGGKKKRYRILKHLVWASRELERFALNPGLLETSEGCKQILGQLQPA
LQTGSEELRSLYNTVATLYCVHQRIEVKDTKEALEKIEEEQNKSQQKTQQAADKGNSSKVSONYPIVQN
LOGQMVHQAIISPRTLNAWVKVIEEKAFSPEVIPMFSALESEGATPQDLNTMLNTVGHHQAAMQMLKDTINE
EEAEWDRLHPVHAGPIPQGMREPRGSDIAGTSTLQEQIAWMTSNPLPVGEIYKRWIILGLINKIVRMY
SPVSILDIRQGPKEPFRDYVDRFFKTLRAEQATODVKNWMTDTLLVQNANPDCKTILKALGPATLEEMM
TACQGVGGPGHKARVLAEMSQTNAIAIMMQRGNFKGQRRRIIKCFCNGKGEGHIAARNCRAPRKKGCWKGK
EGHQMKDCTERQANFLGKIMPSNKGPGNFIQSRPEPTAPPAESFGFGEETPSPKQEPKDKEPPLTSK
SLFGNDPLSQ

Fig. 19E